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Figures 16A and 16B illustrate the preparation of a cleavable tag of the present invention.

Figure 17 illustrates the preparation of an intermediate compound useful in the preparation of a cleavable tag of the invention.

In the Claims

Please cancel claims 1-32, 42-51, and 53-61.

REMARKS

Claims 1-61 were previously pending in the application. By this Preliminary Amendment, claims 1-32, 42-51, and 53-61 have been canceled. Accordingly, claims 33-41 and 52 are currently pending.

Sequence Listing

The enclosed electronic and paper copies of the Sequence Listing include no new matter that goes beyond the original application as filed, but are supplied as requested by the Notice to File Missing Parts. Furthermore, the above amendments, which merely direct the insertion of the Sequence Listing and insertion of sequence identifiers, include no matter that goes beyond the original application as filed. Applicants respectfully submit that the above-identified application is now in compliance with 37 C.F.R. §§ 1.821-1.825 and WIPO Standard ST. 25.

Drawings

Included with this Response is a full set of Formal Drawings which incorporate the changes required by the Notice to File Missing Parts.

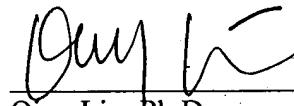
In addition, the amendment also adds a CROSS REFERENCE TO RELATED APPLICATIONS section. No new matter has been added.

Attached hereto is a marked-up version of the changes made to the specification and

claims by the current amendment. The attached page is captioned "**Version With Markings to Show Changes Made.**" Also enclosed is a copy of Limited Recognition Under 37 CFR § 10.9(b).

Consideration of the present application in view of the above amendment is respectfully requested.

Respectfully submitted,
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Seed Intellectual Property Law Group PLLC



Qing Lin, Ph.D.
(See limited recognition)

QXL:jab

Enclosures:

- Copy of Limited Recognition Under 37 CFR § 10.9(b).
- Sequence Listing, Diskette and Declaration
- Filing Formal Drawings, 36 Sheets of Drawings (Figs 1A-17)

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Docket No. : 780068.418C3

VERSION WITH MARKINGS TO SHOW CHANGES MADE

In the Specification:

Paragraph beginning at page 112, line 10, has been amended as follows:

Preparation of 2,4,6-trichlorotriazine derived oligonucleotides: 1000 µg of 5'-terminal amine linked oligonucleotide (5'-hexylamine-TGTAAAACGACGGCCAGT-3") (~~Seq. ID No. 1~~) (SEQ ID NO: 1) are reacted with an excess recrystallized cyanuric chloride in 10% n-methyl-pyrrolidone alkaline (pH 8.3 to 8.5 preferably) buffer at 19 to 25- C for 30 to 120 minutes. The final reaction conditions consist of 0.15 M sodium borate at pH 8.3, 2 mg/ml recrystallized cyanuric chloride and 500 ug/ml respective oligonucleotide. The unreacted cyanuric chloride is removed by size exclusion chromatography on a G-50 Sephadex column.

Paragraph beginning at page 124, line 27, has been amended as follows:

ODNs complementary (5'-CCTTAGGACAGTCTTCACGC; SEQ ID NO: 2) to conserved or hypervariable regions of the 16S ribosomal RNA (rRNA) of *Porphyromonas gingivalis* (Pg), were synthesized on either an ABI 380B or a MilliGen 7500 automated DNA synthesizer using the standard cyanoethyl-N,N-diisopropylamino-phosphoramidite (CED-phosphoramidite) chemistry. Amine tails were incorporated onto the 5'-end using the commercially available N-monomethoxytritylaminohex-6-yloxy-CED-phosphoramidite. ODNs with 5'-monomethoxytrityl groups were chromatographed by HPLC using a Hamilton PRP-1 (7.0 x 305 mm) reversed-phase column employing a gradient of 5% to 45% CH₃CN in 0.1 M

Et₃NH+OAc-, pH 7.5, over 20 min. After detritylation with 80% acetic acid, the ODN s were precipitated by addition of 3 M sodium acetate and 1-butanol. Analytical checks for the quality of the ODNs were done by ion-exchange HPLC using a Toso-Haas DEAE-NPR column and by denaturing polyacrylamide gel electrophoresis (PAGE).

Paragraph beginning at page 127, line 1, has been amended as follows:

F. Solid Support cDNA Synthesis for Gene Expression Assay

Oligo DMO 596 5'- ACTACTGATCAGGCGCGCCTTTTTTTTTTTTTT -3' (SEQ ID NO: 3) spacer Asc I (poly dT)20

Paragraph beginning at page 127, line 24, has been amended as follows:

One nylon bead bearing the covalently linked oligonucleotide, 5'- ACTACTGATCAGGCGCGCCTTTTTTTTTTTT - 3' (SEQ ID NO: 4) (GenSet, La Jolla, CA), is added to 10,ug total cellular RNA, diluted in enough RNase-free water to cover the bead, in a sterile 1.5 ml microfuge tube (Fisher Scientific). The RNA and bead are incubated at 65C for 5 minutes. An equal volume of 2X mRNA hybridization buffer consisting of 50,mM Tris pH 7.5 , 1M NaCl (Fisher Scientific) and 20,ug/ml acetylated-BSA (New England Biolabs, Beverly, MA) is added to each tube and the tubes rocked gently for 2 hours at room temperature. The supernatant is removed and the bead is then washed three times in 1X mRNA hybridization buffer. After the final wash is complete, a reverse transcription mix consisting of 1X MMLV-reverse transcriptase buffer, 1,mM dNTP mix, 2,mM DTT (Life Technologies), 20 units Rnasin (Promega. Madison, WI)and 10,ug/ml acetylated-BS (New England Biolabs) is added to each tube followed by addition of 600 units MMLV-reverse transcriptase(Life Technologies). This reaction is rocked gently at 42°C for 2 hours. 1 unit RNase H (Boehringer-Mannheim. Indianapolis, IN) is then added and the reaction allowed to continue for another 0.5 hour. The supernatant is again removed and each bead is washed three times in 10 mM Tris pH 8.0, 1 mM EDTA pH 8(Fisher Scientific). Remaining RNA template is removed by boiling the beads in TE with 0.01% SDS (Fisher Scientific).

Paragraph beginning at page 128, line 8, has been amended as follows:

The nylon solid support was then hybridized with 100 nanograms per ml of the following tagged oligonucleotide probes

(5'-GAACTCAAACCTCTGGAGGAAGTG-3', IL-2,	<u>(SEQ ID NO: 5)</u>
5'-CAGTGCAGAGGCTCGCGAGCTATA-3', IFN-gamma	<u>(SEQ ID NO: 6)</u>
5'-CTTGACCATGATGGCCAGCCACTA-3', GM-CSF	<u>(SEQ ID NO: 7)</u>
5'-CATTCCCACGGTCACTGCCATCTC-3', c-fos	<u>(SEQ ID NO: 8)</u>
5'-GCGACTGTGCTCCGGCAGTTCTAC-3', IL-4	<u>(SEQ ID NO: 9)</u>
5'-GTGGTTCATCGACGATGCCACGAA-3', PKC-gamma	<u>(SEQ ID NO: 10)</u>
5'-GAGCTCATGTACCCACCTCCGTAC-3', CTLA4/CD28	<u>(SEQ ID NO: 11)</u>
5'-ATCTTGTGCAGCCGCCCTCACTG-3', GMP kinase)	<u>(SEQ ID NO: 12)</u>

Paragraph beginning at page 129, line 19, has been amended as follows:

A "target" oligonucleotide (DMO501: 5'-TTGATTCCAATTATGCGAAGGAG-
3'; SEQ ID NO: 13) was immobilized on a set of solid supports. ODN-beads (3/32nd inch diameter) were prepared as previously described (Van Ness et al., *Nucl. Acids Res.* 19:3345, 1991). The ODN-beads contained 0.01 to 1.2 mg/bead of covalently immobilized ODN. DMO578 is the complement to DMO501 (perfect complement). DMO1969 is the complement to DMO501 with a G--->T change at position 11. DMO1971 is the complement to DMO501 with a A--->T change at position 12. Each probe oligonucleotide was labeled with either BIODIPY, TAMRA or Texas Red. Hybridization reactions were assembled in 3 M GuSCN, 0.01 M Tris pH 7.6, 5 mM EDTA at 50 ng/ml respective probe. Equal molar ratios of each probe type were used in each hybridization in the presence of 3 solid supports per tube. Hybridizations are performed at 42°C for 30 minutes with constant agitation. The beads were washed twice with 3 M GuSCN at 42°C and then with SDS/FW 5 times.

The Section beginning at page 5, line 14 to page 6, line 9, has been revised to read as follows:

BRIEF DESCRIPTION OF THE DRAWINGS

Figures 1A, 1B, and 1C depicts the flowchart for the synthesis of pentafluorophenyl esters of chemically cleavable mass spectroscopy tags, to liberate tags with carboxyl amide termini.

Figures 2A, 2B, and 2C depicts the flowchart for the synthesis of pentafluorophenyl esters of chemically cleavable mass spectroscopy tags, to liberate tags with carboxyl acid termini.

Figures 3-63A, 3B, and 3C; 4A, 4B, and 4C; 5A, 5B, and 5C; 6A, 6B; and 6C; and 8A, 8B, and 8C depict the flowchart for the synthesis of tetrafluorophenyl esters of a set of 36 photochemically cleavable mass spectroscopy tags.

Figures 7A, 7B, and 7C depicts the flowchart for the synthesis of a set of 36 amine-terminated photochemically cleavable mass spectroscopy tags.

Figure 9 depicts the synthesis of 36 photochemically cleavable mass spectroscopy tagged oligonucleotides made from the corresponding set of 36 tetrafluorophenyl esters of photochemically cleavable mass spectroscopy tag acids.

Figures 10A and 10B depicts the synthesis of 36 photochemically cleavable mass spectroscopy tagged oligonucleotides made from the corresponding set of 36 amine-terminated photochemically cleavable mass spectroscopy tags.

Figure 11 illustrates the simultaneous detection of multiple tags by mass spectrometry.

Figure 12 shows the mass spectrogram of the alpha-cyano matrix alone.

Figure 13 depicts a modularly-constructed tagged nucleic acid fragment.

Figure 14 is a schematic representation of an array interrogation system using Matrix Assisted Laser Desorption Ionization (MALDI) mass spectroscopy in accordance with an embodiment of the present invention.

Figures 15A and 15B illustrate the preparation of a cleavable tag of the present invention.

Figures 16A and 16B illustrate the preparation of a cleavable tag of the present invention.

Figure 17 illustrates the preparation of an intermediate compound useful in the preparation of a cleavable tag of the invention.